

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization  
International Bureau



(43) International Publication Date  
2 October 2003 (02.10.2003)

PCT

(10) International Publication Number  
**WO 03/080566 A2**

(51) International Patent Classification<sup>7</sup>: C07C 259/06,  
C07D 207/46, C07C 59/347, C07K 5/06, C07C 309/17,  
C07F 5/02, A61K 31/194, 31/16, 31/33

Peter, John [GB/GB]; Wellcome Trust Centre for Human Genetics, Roosevelt Drive, Oxford OX3 7BN (GB).

(21) International Application Number: PCT/GB03/01239

(74) Agents: ELLIS-JONES, Patrick, George, Armine et al.; J.A. Kemp & Co., 14 South Square, Gray's Inn, London WC1R 5JJ (GB).

(22) International Filing Date: 21 March 2003 (21.03.2003)

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.

(25) Filing Language: English

(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

(26) Publication Language: English

Published:

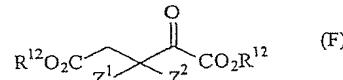
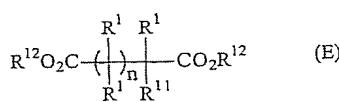
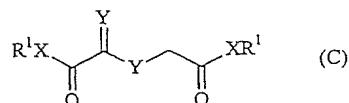
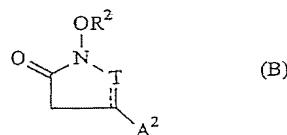
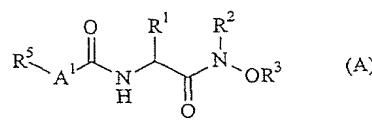
— without international search report and to be republished upon receipt of that report

(71) Applicant (*for all designated States except US*): ISIS INNOVATION LIMITED [GB/GB]; Ewert House, Ewert Place, Summertown, Oxford OX2 7SG (GB).

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(72) Inventors; and  
(75) Inventors/Applicants (*for US only*): SCHOFIELD, Christopher, Joseph [GB/GB]; Dyson Perrins Laboratory, University of Oxford, South Parks Road, Oxford OX1 3QY (GB). MAXWELL, Patrick, Henry [GB/GB]; Imperial College of Science, Technology & Medicine, Hammersmith Campus, Du Cane Road, London W12 0NN (GB). PUGH, Christopher, William [GB/GB]; University of Oxford, Wellcome Trust Centre for Human Genetics, Roosevelt Drive, Oxford OX3 7BN (GB). RATCLIFFE,

(54) Title: HIF HYDROXYLASE INHIBITORS



**WO 03/080566 A2**

(57) Abstract: The invention provides a compound of one of the formulae (A), (B), (C), (D), (E), (F) as herein defined, or a salt thereof, for use in the treatment of a condition associated with increased or decreased HIF levels or activity, or a condition in which an increase or decrease in HIF levels or activity may be beneficial.

## HIF HYDROXYLASE INHIBITORS

### Field of Invention

The present invention relates to compounds which modulate 2OG (2-oxoglutarate) dependent oxygenases, in particular prolyl hydroxylases. These may be useful as modulators of HIF (hypoxia inducible factor) alpha (HIF- $\alpha$ ) prolyl hydroxylase.

### Background of Invention

The transcription factor HIF (hypoxia inducible factor) system is a key regulator of responses to hypoxia, occupying a central position in oxygen homeostasis in a wide range of organisms. A large number of transcriptional targets have been identified, with critical roles in angiogenesis, erythropoiesis, energy metabolism, inflammation, vasomotor function, and apoptotic/proliferative responses. The system is essential for normal development, and plays a key role in pathophysiological responses to ischaemia/hypoxia. HIF is also important in cancer, in which it is commonly upregulated, and has major effects on tumour growth and angiogenesis. The HIF DNA binding complex consists of a heterodimer of  $\alpha$  and  $\beta$  subunits. Regulation by oxygen occurs through hydroxylation of the  $\alpha$ -subunits, which are rapidly destroyed by the proteasome in oxygenated cells. This involves binding of HIF- $\alpha$ -subunits by the von Hippel-Lindau tumour suppressor protein (pVHL), with pVHL acting as the, or part of the, recognition component for a ubiquitin ligase that promotes ubiquitin dependent proteolysis through interaction with a specific sequence or sequences in HIF- $\alpha$ -subunits. In hypoxia, this process is suppressed, so stabilizing HIF- $\alpha$  and permitting transcriptional activation via the HIF  $\alpha, \beta$ .

### Disclosure of the Invention

In our British Application No. 0118952.1 we disclose a polypeptide comprising:

- (a) the amino acid sequence of SEQ ID NO: 2, 4 or 6 having HIF hydroxylase activity;
- 30 (b) a variant thereof having at least 60% identity to the amino acid

sequence of SEQ ID NO: 2, 4 or 6 and having hydroxylase activity; or

(c) a fragment of either thereof having HIF- $\alpha$  hydroxylase activity.

Preferably, the polypeptides have prolyl hydroxylase activity and require Fe(II) for activity.

5 They are related by sequence to non-haem oxygenases for which crystal structures are known, e.g. proline-3-hydroxylase (Clifton et al, Eur. J. Biochem, 2001, 268, 6625-6636).

It also discloses polynucleotides which encode the polypeptides as well as expression vectors comprising the polynucleotide and antibodies capable of 10 specifically binding the polypeptide. We also disclose assays for identifying modulators of the activity of the HIF hydroxylase as well as the use of modulators such as inhibitors of the activity of the peptides in the treatment of a condition or disease associated with altered HIF levels with respect to healthy (or normal) levels, and the treatment of conditions where an alteration in the HIF levels or activity 15 would be beneficial.

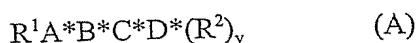
Inhibitors of the 2-OG dependent enzyme collagen prolyl-4-hydroxylase (CPH) are well known in the art and have been previously proposed for use in the treatment of lung fibrosis, skin fibrosis (scleroderma), atherosclerosis and other conditions associated with collagen biosynthesis. Inhibitors of para-hydroxyphenylpyruvate oxygenase (a non-haem oxygenase employing ferrous iron as a co-factor) such as triketones are used as herbicides (Lee D. et al (1998) Pestic. Sci. 54(4) 377-384). We have disclosed that certain of these CPH inhibitors (and other components) also inhibit the biological (i.e. HPH) activity of an PHD polypeptide. A CPH inhibitor or modified CPH inhibitor which inhibits the biological activity of an 20 PHD polypeptide may be used in the treatment of a condition associated with reduced or suboptimal HIF levels or activity, or a condition in which an increase in HIF levels or activity may be beneficial, for example ischaemia, wound healing, auto-, allo-, and xeno- transplantation, systemic high blood pressure, cancer, inflammatory disorders, and metabolic disorders.

30 Various methods and uses of modulators which inhibit, potentiate, increase or stimulate hydroxylation of HIF- $\alpha$  by an PHD polypeptide are disclosed. The purpose

of disruption, interference with or modulation of the hydroxylation of HIF-1 $\alpha$  by a PHD polypeptide may be to modulate cellular functions such as angiogenesis, erythropoiesis, energy metabolism, inflammation, matrix metabolism, vasomotor function, and apoptotic/proliferative responses and pathophysiological responses to ischaemia/hypoxia, all of which are mediated by HIF $\alpha$  as discussed above.

Compounds which modulate 2OG oxygenases, in particular CPH may be useful as modulators of HIF prolyl hydroxylase, or may be used as 'lead' compounds which may be modified and/or optimised to develop modulators of HIF prolyl hydroxylase, in particular selective modulators are described.

Some of these compounds generally possess the formula:



where the group R<sup>1</sup> is capable of forming an electrostatic interaction with the sidechain of the arginine residue which, together with other residues, binds the 5-carboxylate of 2-oxoglutarate during catalysis, A\*B is a chain of two atoms which are, independently, carbon, oxygen, nitrogen or sulphur, which chain can be functionalised, y is 0 or 1 and C\*D is a chain of two atoms which are, independently, carbon, oxygen, nitrogen, or sulphur, which chain can be functionalised, A, B, C and D being linked to one another by a single and/or double and/or triple bond such that when y is 0 or 1 at least one of the atoms of which is capable of chelating with a metal group and when y is 1 said chain is attached to R<sup>2</sup> which is capable of chelating with a metal group. Generally at least one of A, B, C and D is not carbon. Typical chains include C-N-C-C and C-O-C-C and C-C-C=O. The chain atoms can form part of a ring.

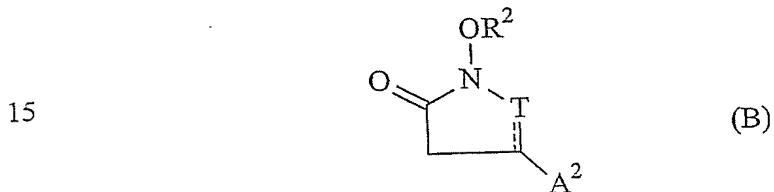
New classes of modulators of HIF prolyl hydroxylase have been found, according to the present invention. These possess the following formulae (A) to (F)



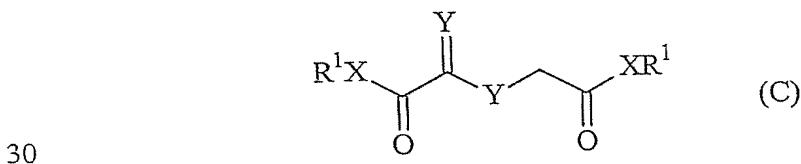
30

where each of R<sup>1</sup> and R<sup>5</sup> is independently H, OH, SH, a branched or straight C<sub>1</sub> to C<sub>6</sub>

alkyl chain optionally containing 1 or more eg. 2 N, S, O or P chain atoms, especially methyl, which can be functionalised, any amino acid side chain, such as alanine, phenylalanine, valine and glutamic acid, a 4 to 7 membered heterocyclic ring 5 optionally containing 1 or 2 N, S, O or P atoms or a 5 or 6 membered aromatic ring, optionally containing 1 or 2 N, O or S atoms which may be fused to another ring or a said alkyl chain substituted by a said aromatic ring, such as aryloxy alkyl, A<sup>1</sup> is CH<sub>2</sub> or O, and each of R<sup>2</sup> and R<sup>3</sup> is independently be H, OH, a branched or straight C<sub>1</sub> to C<sub>6</sub> alkyl chain optionally containing 1 or more eg. 2 N, S, O or P chain atoms which 10 can be functionalised, optionally with 1, 2, 3, 4 or 5 halo substitutions, a 4 to 7 membered heterocyclic ring optionally containing 1 or 2 N, S, O or P atoms, or a 5 or 15 6 membered aromatic ring, optionally containing 1 or 2 N, O or S atoms which may be fused to another ring or a said alkyl chain substituted by a said aromatic ring,



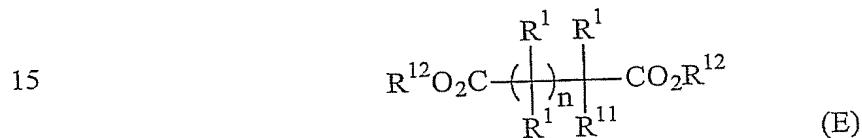
wherein R<sup>2</sup> is as defined above, - - - is a single bond and T is CH<sub>2</sub> or C=O, or - - - is a double bond and T is CH; A<sup>2</sup> is H or -XCO<sub>2</sub>R<sup>4</sup>; X is a single bond or a branched or 20 straight C<sub>1</sub> to C<sub>6</sub> alkyl chain, optionally containing 1 or more eg. 2 N, S, O or P chain atoms and optionally substituted by eg. halo, OH, NHR<sup>2</sup> or NHCOR<sup>4</sup> where R<sup>2</sup> and R<sup>4</sup> are as defined above and R<sup>4</sup> represents H, a branched or straight chain C<sub>1</sub> to C<sub>6</sub> alkyl group optionally containing 1 or more eg. 2 N, S, O or P chain atoms, a 4 to 7 membered heterocyclic ring, optionally containing 1 or 2 N, S, O or P atoms, or a 5 25 membered aromatic ring, optionally containing 1 or 2 N, O or S atoms, which may be fused to another ring, or a salt thereof,



where each X which may be the same or different is NH, NR", where R" is OH, a branched or straight C<sub>1</sub> to C<sub>6</sub> alkyl chain optionally containing 1 or more eg. 2 N, S, O or P chain atoms which can be functionalised, or O i.e. XR<sup>1</sup> is typically OH or O-alkyl having a branched or straight C<sub>1</sub> to C<sub>6</sub> alkyl chain, especially MeO,  
5 each Y, which may be the same or different, is O or S and each R<sup>1</sup>, which may be the same or different, is as defined above,



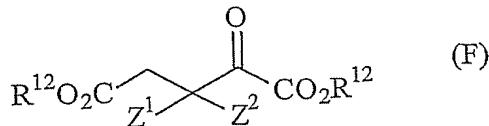
10 where m is 0 or 1, Q represents (R<sup>1</sup>R<sup>6</sup>)<sub>x</sub>Z where x is 0, 1 or 2, R<sup>1</sup> is as defined above and R<sup>6</sup> is as defined for R<sup>1</sup>, and Z is P=O(OH)<sub>2</sub>, B(OH)<sub>2</sub> or SO<sub>3</sub>H, or a salt thereof, typically a sodium salt, or



where each R<sup>1</sup>, which are the same or different, is as defined above; R<sup>11</sup> represents OH or R<sup>10</sup> NH where R<sup>10</sup> is HO, R<sup>1</sup>CO or HOOC(X)<sub>x</sub> wherein R<sup>1</sup> is as defined above,  
20 x is 0 or 1 and X is R<sup>1</sup>R<sup>1</sup>C wherein each R<sup>1</sup>, which are the same or different, is as defined above; or R<sup>10</sup> is an amino acid residue H<sub>2</sub>N (R<sup>1</sup>R<sup>1</sup>C) CO - wherein each R<sup>1</sup>, which are the same or different, is as defined above; n is 1 or 2 and R<sup>12</sup> is H or straight or branched C<sub>1</sub> to C<sub>6</sub> alkyl; or a salt thereof. Typically X is CH<sub>2</sub> or CHOH.

Another aspect of the invention concerns analogues of 2-oxoglutarate that act as improved (relative to 2-oxoglutarate) co-substrates for the HIF hydroxylases.  
25 Such a compound is 3-fluoro 2-oxoglutarate. Assays in which this compound replaces 2-oxoglutarate demonstrate a higher level of HIF hydroxylation than observed when using 2-oxoglutarate under analogous conditions.

These analogues possess the formula:



5       wherein each of  $Z^1$  and  $Z^2$  is independently hydrogen, SH or an electron withdrawing group such as halogen, preferably fluorine, or alkoxy such as methoxy, and  $R^{12}$  is as defined above, or a salt thereof. Preferably one of  $Z^1$  and  $Z^2$  is hydrogen and the other is fluorine (3-F-2-OG).

Accordingly the present invention provides a compound of formula (A) to  
10     (F) for use in the treatment of a condition associated with increased or decreased HIF levels or activity, or a condition in which an increase or decrease in HIF levels or activity may be beneficial, as well as the use of a compound of formula (A) to (F) in the manufacture of a medicament for the treatment of such a condition.

The said alkyl groups and chains are typically functionalised by alcohol,  
15     fluorine, thiol, a carboxylic acid, phosphonic or phosphinic acid, sulphonate acid or other chelating group, in the case of the chains typically via an alkyl group.

In the formulae described herein, a branched or straight  $C_1$  to  $C_6$  alkyl chain may be a methyl, ethyl, propyl, butyl, iso-butyl, *tert*-butyl, pentyl, neopentyl, *tert*-pentyl or a primary, secondary or tertiary hexyl group. Hetero atoms such as O, S, N and P may replace one or more of the carbon atoms. Preferably the alkyl groups are methyl, the preferred heterocyclic rings are pyrrolidine and tetrahydropyran and the preferred aromatic rings are benzene, naphthalene and pyridine.

Typically, each of  $R^1$  and  $R^5$  is independently H, OH, a branched or straight  $C_1$  to  $C_6$  alkyl chain optionally containing 1 or more N, S, O or P chain atoms, which  
25     can be functionalised, any amino acid side chain, a 4 to 7 membered heterocyclic ring optionally containing 1 or 2 N, S, O or P atoms or a 5 or 6 membered aromatic ring, optionally containing 1 or 2 N, O or S atoms which may be fused to another ring or a said alkyl chain substituted by a said aromatic ring.

Typically,  $A^1$  is  $CH_2$ .

30     Typically,  $A^2$  is  $-XCO_2R^4$ .

Typically,  $R^{11}$  represents  $R^{10}NH$  where  $R^{10}$  is  $R^1CO$  or  $HOOC(X)_x$  wherein

$R^1$  is as defined above,  $x$  is 0 or 1 and  $X$  is  $R^1R^1C$  wherein each  $R^1$ , which are the same or different, is as defined above; or  $R^{10}$  is an amino acid residue  $H_2N(R^1R^1C)CO-$  wherein each  $R^1$ , which are the same or different, is as defined above.

Typically, each of  $Z^1$  and  $Z^2$  is independently hydrogen or an electron withdrawing group.

Typically, in the compounds of formula (F),  $R^{12}$  is H. Alternatively,  $R^{12}$  may be straight or branched  $C_1$  to  $C_6$  alkyl.

The compounds of formula (A) are hydroxamates. Preferred compounds include those where  $R^5$  is aryloxyalkyl, especially oxyloxyethyl such as phenyloxymethyl or phenylalkyloxymethyl, especially benzyloxymethyl or substituted benzyloxymethyl such as p-hydroxy benzyloxymethyl and/or where  $R^2$  and/or  $R^3$  is  $HOCH_2$ .

Typical compounds include N-phenoxy-acetyl-(L)-alanine-hydroxamide (Is41) and the corresponding (D) isomer (Is43) as well as the corresponding tyrosine derivatives (Is44 and 45) and L- and D-phenylglycine derivatives (Is46 and 47), along with benzo hydroxamic acid and N-phenoxyacetyl-D-phenylalanine hydroxamic acid (Is42).

These compounds can generally be prepared following the method of Walter et al., Tetrahedron 1997, 53, 7275-7290 and Biorg. Chem 1999, 27, 35-40.

The compounds of formula (B) are cyclic hydroxamates. Preferred compounds are those where  $X$  is a single bond or methyl and/or  $R^2$  is H or phenylalkyl, especially benzyl and/or  $R^4$  is H or methyl. Typical compounds include (1-hydroxy-2, 5-dioxo-pyrrolidin-3-yl) acetic acid (Is52), (1-hydroxy-2, 5-dioxo-pyrrolidin-3-yl) carboxylic acid (ANU 2) and its N-benzyloxy derivative (ANU 1) along with (1-benzyloxy-2, 5-dioxo-pyrrolidin-3-yl) acetic acid (Is50) and the corresponding methyl ester (Is64), and N-hydroxy succinimide (C1). Note that Is52 ( $R^2=H$ ,  $T=C=O$ ,  $X=CH_2$ ,  $R^4=H$ ) is highly active reflecting its structural analogy with 2-oxoglutarate. These compounds can be prepared using the general procedure of Schlicht et al. (US 4446038).

The compounds of formula (C) are analogues of 2-oxoglutarate or oxalyl derivatives of hydroxyacetate and mercapto acetic acid. Preferred compounds

include those where X is O and/or R<sup>1</sup> is H or methyl. Typical compounds include dimethyl oxalylglycolate (Is10) as well as its free acid (Is14) and dimethyl oxalylthioglycolate (Is11). These compounds can be prepared following Franklin et al., J. Med. Chem 1992, 35, 2652-2658 or Kwon et al., J. Am. Chem. Soc. 1989, 111,

5 1854-1860.

The compounds of formula (D) are carboxylic acids which possess a phosphonic, sulphonic or boronic acid group as well as salts of these. Typically R<sup>1</sup> and R<sup>6</sup> are hydrogen. Preferred compounds include the phosphoric acids where x is 0, 1 or 2 (C3, 4 and 5, respectively) as well as disodium 3-sulpho-propionate (Is63) 10 and its free acid, and 3-borono-propionic acid (Is62).

The compounds of formula (E) are N-acylated amino acids or polycarboxylic acids. Typical compounds are those where R<sup>1</sup> is H, and/or R<sup>12</sup> is H or ethyl. When R<sup>11</sup> represents R<sup>10</sup>NH the compounds are typically dipeptides such that R<sup>10</sup> is an acyl group of a natural amino acid such as glycine. Typical preferred such compounds 15 include Asp-Gly (C18), cyclo (Asp-Gly) (C19), beta-Asp-Gly (C20), Glu-Gly (C21) and Z-Glu-Gly (C22). Other typical compounds include those where R<sup>10</sup> is acetyl or benzoyl such as the N-acetylated derivatives of L-aspartic acid (C6) and of L-glutamic acid (C7) i.e. R<sup>10</sup> is acetyl and N-benzoylated derivatives of glutamic acid (C15 and Is90) i.e. R<sup>10</sup> is benzoyl. Other typical compounds include those where R<sup>11</sup> 20 is -NHOH such as diethyl 2-(hydroxylamino)-glutarate (Is51 being the racemic form of this compound) and those where R<sup>11</sup> is OH such as 2-hydroxyglutaric acid (Is57). When R<sup>11</sup> is HOOC(X)x, X is especially CH<sup>2</sup> or CHOH. The compounds are typically citric acid (C12), tricarballylic acid (C13) and succinic acid as well as the tri-methyl ester of ethane tricarboxylic acid (Is72).

25 The compounds of formula (F) are analogues of 2-oxoglutarate. Preferred compounds include 3-fluoro-2-oxoglutarate compounds (i.e. Z<sup>1</sup> is H and Z<sup>2</sup> is F) such as 3-fluoro-2-oxoglutaric acid (Is18) and the corresponding dimethyl ester (Is19).

The compounds which are acids can be present in the form of salts, such as sodium salts.

30 For therapeutic treatment, the compound may be used in combination with any other active substance, e.g. for anti-tumour therapy another anti-tumour

compound or therapy, such as radiotherapy or chemotherapy.

Generally, the modulator is provided in an isolated and/or purified form, i.e. substantially pure. This may include being in a composition where it represents at least about 90% active ingredient, more preferably at least about 95%, more 5 preferably at least about 98%. Any such composition may, however, include inert carrier materials or other pharmaceutically and physiologically acceptable excipients, such as those required for correct delivery, release and/or stabilisation of the active agent. As noted below, a composition according to the present invention may include in addition to a modulator compound as disclosed, one or more other molecules of 10 therapeutic use, such as an anti-tumour agent.

In general they take the form of compositions wherein the compound is in a mixture with a pharmaceutically acceptable carrier or diluent. The carrier may be liquid, e.g. saline, ethanol, glycerol and mixtures thereof, or solid, e.g. in the form of a tablet, or in a semi-solid form such as a gel formulated as a depot formulation or in 15 a transdermally administerable vehicle, such as a transdermal patch. The modulator compound or composition comprising it may be formulated as the coating of a coated stent.

The invention further provides a method of treatment which includes administering to a patient compound as defined above. Exemplary purposes of such 20 treatment are discussed elsewhere herein.

The therapeutic/prophylactic purpose of such a method or use may be the modulation of the level of HIF $\alpha$  in a cell by modulation, e.g. disruption or interference, of the hydroxylation of HIF $\alpha$ , which may occur for example at proline 402, 564 or other proline residue. Hydroxylation of HIF $\alpha$  promotes pVHL binding 25 which leads to ubiquitin dependent proteolysis of HIF $\alpha$  as described above.

The therapeutic/prophylactic purpose may be related to the treatment of a condition associated with reduced or suboptimal or increased HIF levels or activity, or conditions where an alteration in HIF levels or activity may be beneficial such as:  
(i) ischaemic conditions, for example organ ischaemia, including coronary,  
30 cerebrovascular and peripheral vascular insufficiency. The therapy may be applied in two ways; following declared tissue damage, e.g. myocardial infarction (in order to

limit tissue damage), or prophylactically to prevent or ameliorate ischaemia, e.g. promotion of coronary collaterals in the treatment of angina.

(ii) wound healing and organ regeneration.

(iii) auto-, allo-, and xeno- transplantation.

5 (iv) systemic blood pressure.

(v) cancer; HIF $\alpha$  is commonly up-regulated in tumour cells and has major effects on tumour growth and angiogenesis.

(vi) inflammatory disorders.

(vii) pulmonary arterial blood pressure, neurodegenerative disease.

10 (viii) metabolic disorders, e.g. diabetes.

Modulating HIF prolyl hydroxylase activity in a person, an organ, or a group of cells may be exploited in different ways to obtain a therapeutic benefit:

(a) Non cell autonomous: The HIF system is used by cells to influence the production of substances which signal to other cells. These signals may then have

15 effects at (i) a distant site (for example erythropoietin acts on the bone marrow) or (ii) locally (angiogenic growth factors increase the local formation of blood vessels).

Manipulating non cell autonomous behaviour via altering hydroxylase activity is therefore useful in the treatment of anaemia, and local ischaemia, for example in the eye, brain, heart and limbs. Many other signals that are involved in aspects of

20 physiological homeostasis may be, or are known to be, adjusted by HIF activation.

Consequently altering HIF prolyl hydroxylase activity may be used to potentiate or initiate a helpful response for a therapeutic benefit, or to prevent or ameliorate a harmful response. For example, this approach can be used to alter appetite, or blood pressure in the systemic or pulmonary beds.

25 (b) Cell autonomous: the HIF system is also used by cells to regulate cellular metabolism, and decisions concerning differentiation, proliferation and apoptosis. Therefore manipulating the HIF system can be used to alter the viability and behaviour of cells. An increase in cell viability can be achieved by increasing HIF activation, for example in an ischaemic tissue. This approach can also be used

30 in improving pancreatic beta cell viability as a way of ameliorating diabetes, or of improving the viability or function of a group or groups of neurons in Parkinson's

disease, motorneurone disease or forms of dementia. In a different approach, the HIF signal can be manipulated to prevent a group of cells proliferating, or to promote its death or differentiation. For example transient activation of the HIF system in a malignant tumour can be used to provoke death of a substantial number of tumour  
5 cells.

#### Pharmaceutical Compositions

In various further aspects, the present invention thus provides a pharmaceutical composition, medicament, drug or other composition for such a purpose, the composition comprising one or more compounds of formulae (A) to (F), or derivatives thereof, the use of such an composition in a method of medical treatment, a method comprising administration of such a composition to a patient, e.g. for treatment (which may include preventative treatment) of a medical condition as described above, use of such an agent compound or substance in the manufacture 10 of a composition, medicament or drug for administration for any such purpose, e.g. for treatment of a condition as described herein, and a method of making a pharmaceutical composition comprising admixing such an agent, compound or substance with a pharmaceutically acceptable excipient, vehicle or carrier, and optionally other ingredients.  
15

The agent may be used as sole active agent or in combination with one another or with any other active substance, e.g. for anti-tumour therapy another anti-tumour compound or therapy, such as radiotherapy or chemotherapy.  
20

Whatever the agent used in a method of medical treatment of the present invention, administration is preferably in a "prophylactically effective amount" or a "therapeutically effective amount" (as the case may be, although prophylaxis may be considered therapy), this being sufficient to show benefit to the individual. The actual amount administered, and rate and time-course of administration, will depend 25 on the nature and severity of what is being treated. Prescription of treatment, e.g. decisions on dosage etc, is within the responsibility of general practitioners and other medical doctors.  
30

An agent or composition may be administered alone or in combination with

other treatments, either simultaneously or sequentially dependent upon the condition to be treated, e.g. as described above.

Pharmaceutical compositions according to the present invention, and for use in accordance with the present invention, may include, in addition to active 5 ingredient, a pharmaceutically acceptable excipient, carrier, buffer, stabiliser or other materials well known to those skilled in the art. Such materials should be non-toxic and should not interfere with the efficacy of the active ingredient. The precise nature of the carrier or other material will depend on the route of administration, which may be oral, or by injection, e.g. cutaneous, subcutaneous or intravenous. The 10 compositions will typically be sterile.

Pharmaceutical compositions for oral administration may be in tablet, capsule, powder or liquid form. A tablet may include a solid carrier such as gelatin or an adjuvant. Liquid pharmaceutical compositions generally include a liquid carrier such as water, petroleum, animal or vegetable oils, mineral oil or synthetic oil. 15 Physiological saline solution, dextrose or other saccharide solution or glycols such as ethylene glycol, propylene glycol or polyethylene glycol may be included.

For intravenous, cutaneous or subcutaneous injection, or injection at the site of affliction, the active ingredient will be in the form of a parenterally acceptable aqueous solution which is pyrogen-free and has suitable pH, isotonicity and stability. 20 Those of relevant skill in the art are well able to prepare suitable solutions using, for example, isotonic vehicles such as Sodium Chloride Injection, Ringer's Injection, Lactated Ringer's Injection. Preservatives, stabilisers, buffers, antioxidants and/or other additives may be included, as required.

Liposomes, particularly cationic liposomes, may be used in carrier 25 formulations. Examples of techniques and protocols mentioned above can be found in Remington's Pharmaceutical Sciences, 16th edition, Osol, A. (ed), 1980.

The substance or composition may be administered in a localised manner to a particular site or may be delivered in a manner in which it targets particular cells or tissues, for example using intra-arterial stent based delivery.

30 Targeting therapies may be used to deliver the active substance more specifically to certain types of cell, by the use of targeting systems such as antibody

or cell specific ligands. Targeting may be desirable for a variety of reasons, for example if the agent is unacceptably toxic, or if it would otherwise require too high a dosage, or if it would not otherwise be able to enter the target cells.

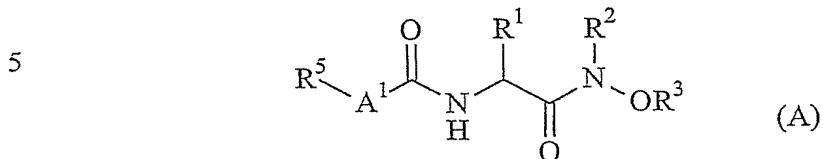
5 The following Examples further illustrate the present invention.

Example 1

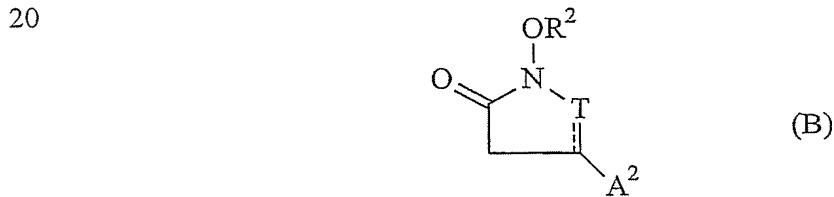
In vitro screening of potential inhibitors of HIF modification was performed using a capture assay. A Gal/HIF-1 $\alpha$ /VP16 fusion protein expressing HIF-1 $\alpha$  residues 549-582 was prepared by IVTT (see British Application No. 0118952.1) and 10 used as a substrate in the assay. The unlabelled substrate was immunopurified on beads, washed, and aliquots incubated in the presence of RCC4 cell extract, with 100 $\mu$ M FeCl<sub>2</sub> and 2mM of the potential inhibitor. The inhibitors were either dissolved in DMSO or Tris as indicated. Controls, where no inhibitor but the equivalent amount of DMSO or Tris was added, were also performed. After washing, 15 the beads were assayed for their ability to interact with 35-S labelled pVHL IVTT. Hydroxylation of the fusion protein by HIF hydroxylase present in the cell extract leads to the ability to capture the labelled pVHL and the amount of labelled protein bound to the fusion protein can then be measured to determine relative HIF hydroxylase activity. Figures 1 to 5 show HIF hydroxylase activity in the presence of 20 a particular inhibitor relative to that seen in the absence of the inhibitor (the DMSO/Tris control).

CLAIMS

1. A compound of one of the formulae

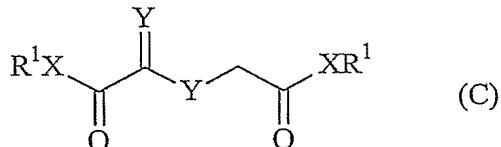


where each of R<sup>1</sup> and R<sup>5</sup> is independently H, OH, SH, a branched or straight C<sub>1</sub> to C<sub>6</sub> alkyl chain optionally containing 1 or more N, S, O or P chain atoms, which can be functionalised, any amino acid side chain, a 4 to 7 membered heterocyclic ring optionally containing 1 or 2 N, S, O or P atoms or a 5 or 6 membered aromatic ring, optionally containing 1 or 2 N, O or S atoms which may be fused to another ring or a said alkyl chain substituted by a said aromatic ring, A<sup>1</sup> is CH<sub>2</sub> or O, and each of R<sup>2</sup> and R<sup>3</sup> is independently H, OH, a branched or straight C<sub>1</sub> to C<sub>6</sub> alkyl chain optionally containing 1 or more N, S, O or P chain atoms which can be functionalised, optionally with 1, 2, 3, 4 or 5 halo substitutions, a 4 to 7 membered heterocyclic ring optionally containing 1 or 2 N, S, O or P atoms, or a 5 or 6 membered aromatic ring, optionally containing 1 or 2 N, O or S atoms which may be fused to another ring or a said alkyl chain substituted by a said aromatic ring,



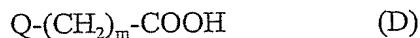
25 wherein R<sup>2</sup> is as defined above, — is a single bond and T is CH<sub>2</sub> or C=O, or — is a double bond and T is CH; A<sup>2</sup> is H or -XCO<sub>2</sub>R<sup>4</sup>; X is a single bond or a branched or straight C<sub>1</sub> to C<sub>6</sub> alkyl chain optionally containing 1 or more N, S, O or P chain atoms, and optionally substituted by halo, OH, OR<sup>4</sup>, NRR<sup>2</sup> or NHCOR<sup>4</sup> where R<sup>2</sup>, R<sup>3</sup> and R<sup>4</sup> are as defined above and R<sup>4</sup> represents H, a branched or straight chain C<sub>1</sub> to C<sub>6</sub> alkyl group, a 4 to 7 membered heterocyclic ring, optionally containing 1 or 2 N, S, O or P atoms, or a 5 or 6 membered aromatic ring, optionally containing 1 or 2 N,

O or S atoms, which may be fused to another ring,

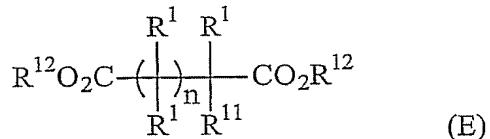


5

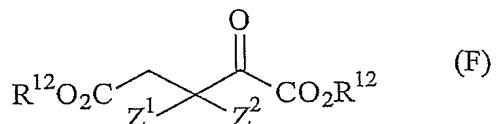
where each X which may be the same or different is NH, NR'', where R'' is OH, a branched or straight C<sub>1</sub> to C<sub>6</sub> alkyl chain optionally containing 1 or more eg. 2 N, S, O or P chain atoms which can be functionalised, or O, each Y, which may be the same or different, is O or S and each R<sup>1</sup>, which may be the same or different, is as defined above,  
10 defined above,



where m is 0 or 1, Q represents (CR<sup>1</sup>R<sup>6</sup>)<sub>x</sub>Z where x is 0, 1 or 2, R<sup>1</sup> is as defined above  
15 and R<sup>6</sup> is as defined for R<sup>1</sup> and Z is P=O(OH)<sub>2</sub>, B(OH)<sub>2</sub> or SO<sub>3</sub>H, or a salt thereof, or



20 where each R<sup>1</sup>, which are the same or different, is as defined above; R<sup>11</sup> represents OH or R<sup>10</sup> NH where R<sup>10</sup> is HO, R<sup>1</sup>CO or HOOC(X)<sub>x</sub> wherein R<sup>1</sup> is as defined above, x is 0 or 1 and X is R<sup>1</sup>R<sup>1</sup>C wherein each R<sup>1</sup>, which are the same or different, is as defined above; or R<sup>10</sup> is an amino acid residue H<sub>2</sub>N (R<sup>1</sup>R<sup>1</sup>C) CO- wherein each R<sup>1</sup>, which are the same or different, is as defined above; n is 1 or 2 and R<sup>12</sup> is H or  
25 straight or branched C<sub>1</sub> to C<sub>6</sub> alkyl; or a salt thereof,



30 wherein each of Z<sup>1</sup> and Z<sup>2</sup> is independently hydrogen, SH or an electron withdrawing group, and R<sup>12</sup> is as defined above, or a salt thereof for use in the treatment of a

condition associated with increased or decreased HIF levels or activity, or a condition in which an increase or decrease in HIF levels or activity may be beneficial.

2. A compound according to claim 1 which is of formula (A) where R<sup>1</sup> is aryloxymethyl, R<sup>2</sup> is hydrogen and R<sup>3</sup> is hydrogen.

5 3. A compound according to claim 2 wherein R<sup>1</sup> is phenoxyxymethyl, benzyloxyxymethyl or p-hydroxybenzylmethyl.

4. A compound according to claim 1 of formula (B) where X is a single bond or methyl, R<sup>2</sup> is H or phenyl alkyl and R<sup>4</sup> is H or methyl.

10 5. A compound according to claim 1 of formula (C) wherein X is O and R<sup>1</sup> is H or methyl.

6. A compound according to claim 1 of formula (D) wherein x is 1, and R<sup>1</sup> and R<sup>6</sup> are both hydrogen.

7. A compound according to claim 1 of formula (E) wherein R<sup>1</sup> is H and R<sup>12</sup> is H or ethyl.

15 8. A compound according to claim 7 wherein R<sup>11</sup> represents R<sup>10</sup> NH and R<sup>10</sup> is an acyl group of a natural amino acid, acetyl or benzoyl.

9. A compound according to claim 7 wherein R<sup>11</sup> is HOOC(X)x.

10. Use of a compound as defined in claim 1 in the manufacture of a medicament for the treatment of a condition associated with increased or decreased HIF levels or activity, or a condition in which an increase or decrease in HIF levels or activity may be beneficial.

20 11. Use according to claim 10 wherein the compound is as defined in any one of claims 2 to 9.

12. A pharmaceutical composition which comprises a compound as defined in 25 claim 1 together with a pharmaceutically acceptable excipient vehicle or carrier.

13. A composition according to claim 12 wherein the compound is as defined in any one of claims 2 to 9.

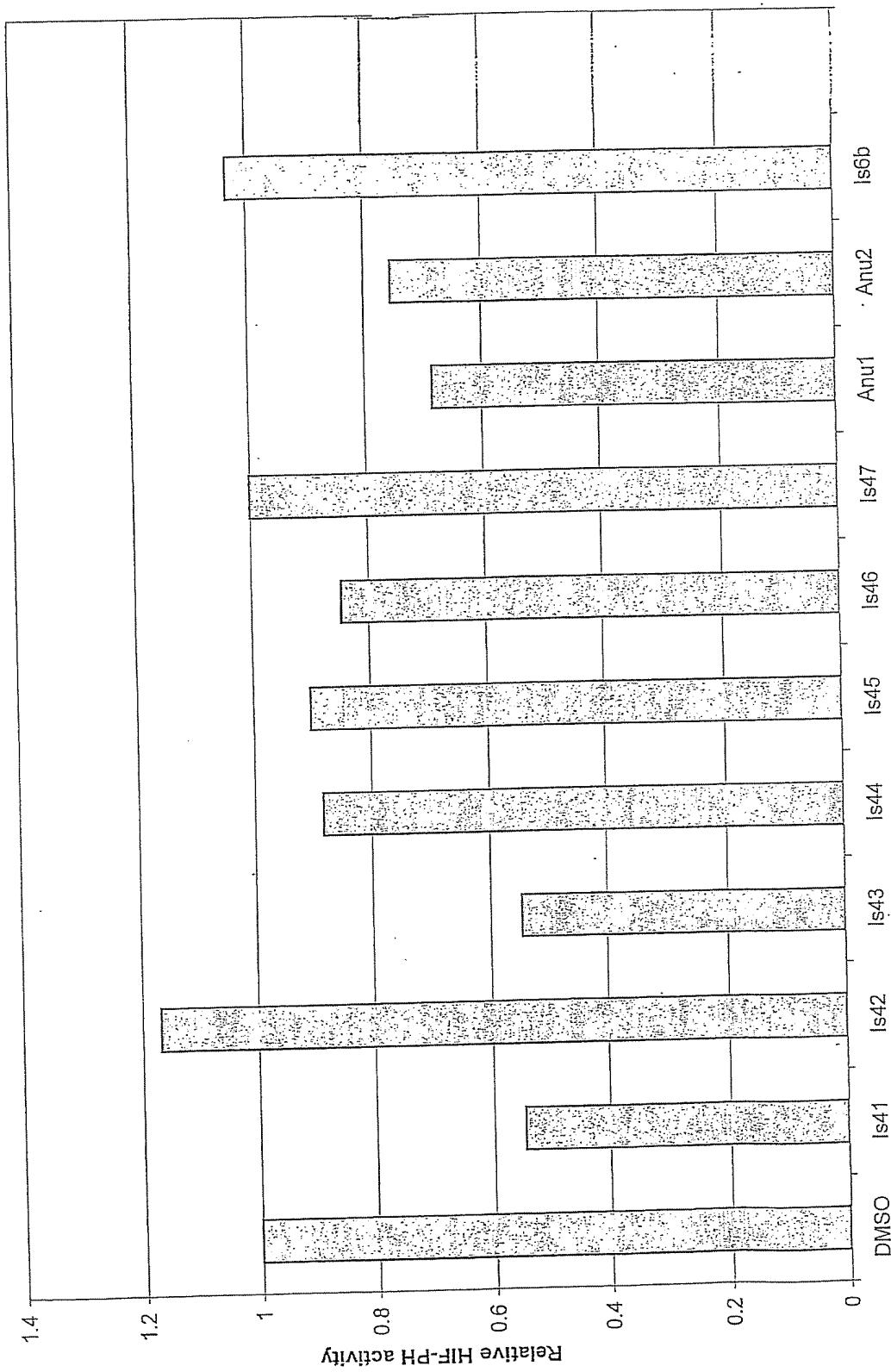
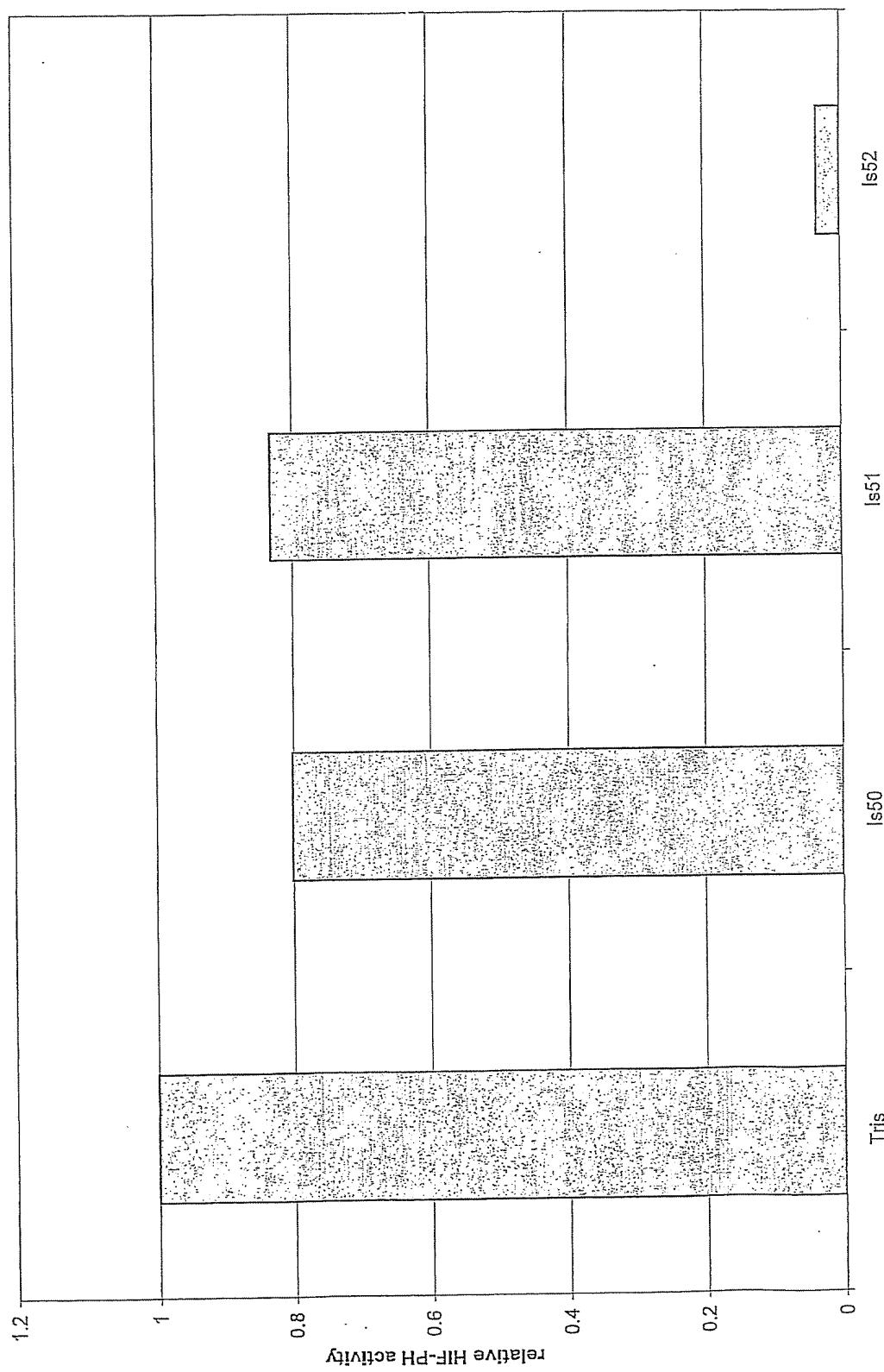
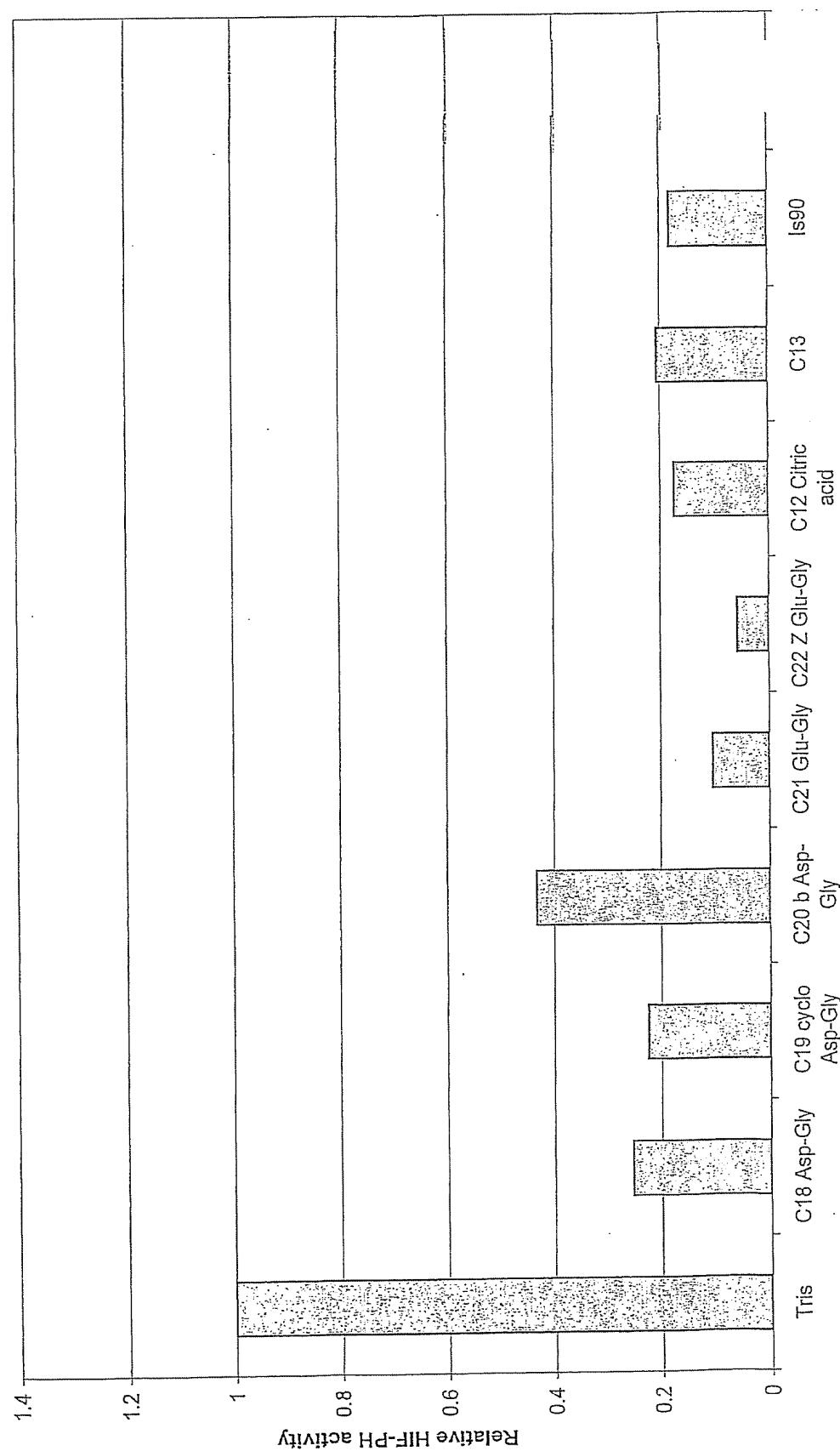
**Figure 1**

Figure 2



**Figure 3**

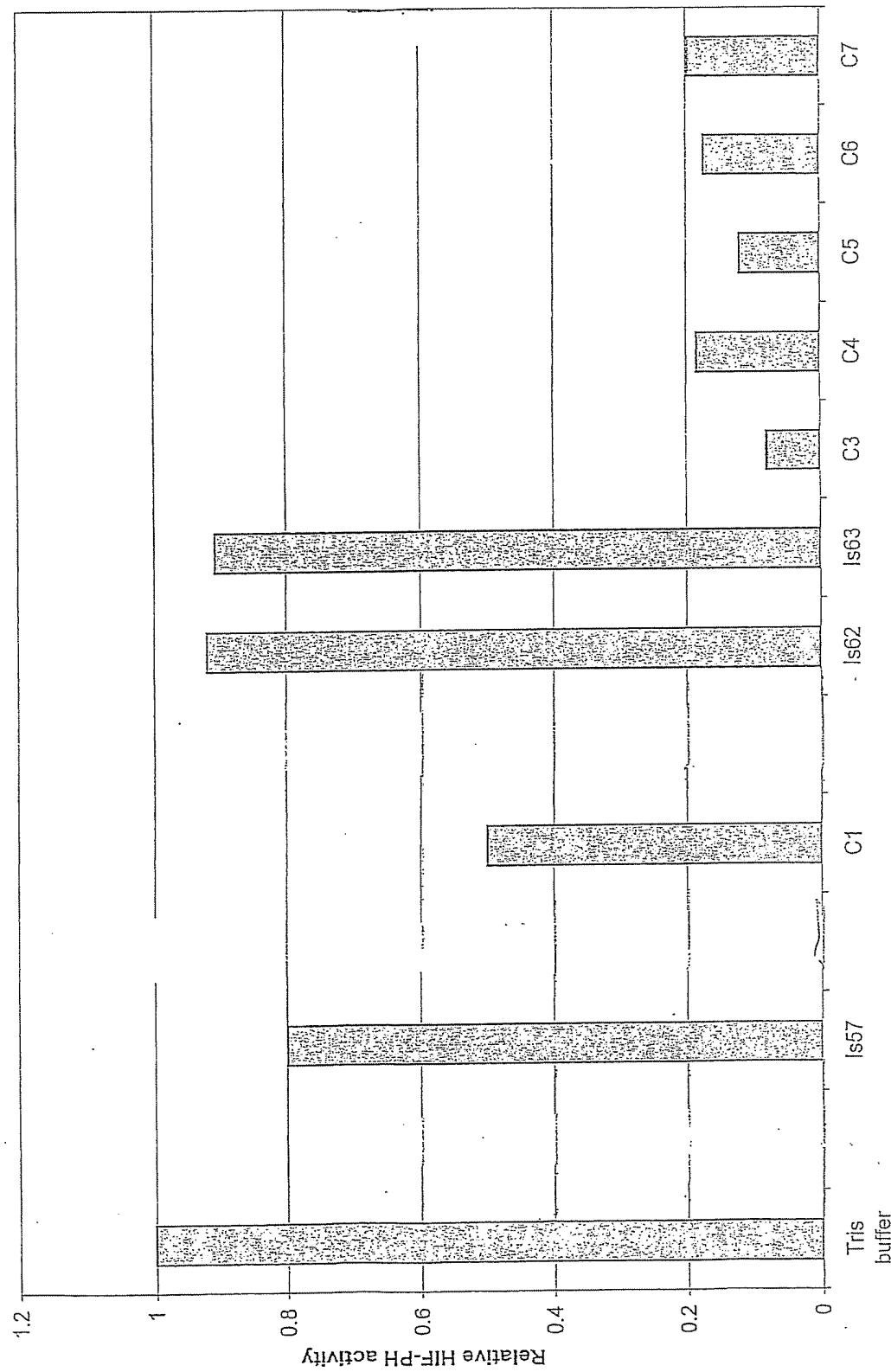
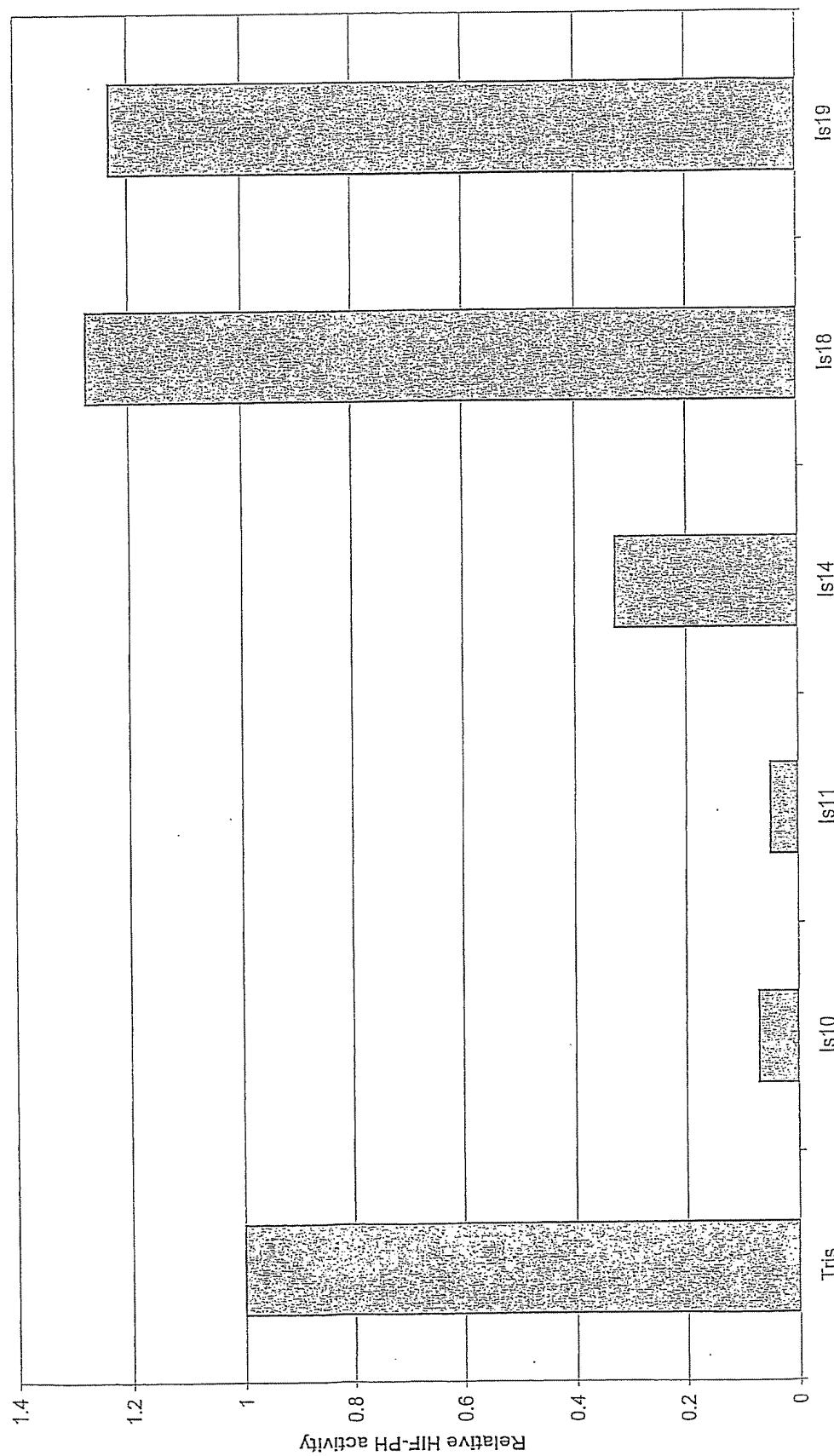
**Figure 4.**

Figure 5



## SEQUENCE LISTING

<110> ISIS INNOVATION LIMITED  
 <120> HIF HYDROXYLASE INHIBITORS  
 5 <130> N.85209A PEJ  
  
 <160> 6  
 <170> PatentIn version 3.1  
  
 10 <210> 1  
 <211> 2110  
 <212> DNA  
 <213> Homo sapiens  
 <220>  
 15 <221> CDS  
 <222> (297)..(1517)  
 <223>  
  
 <400> 1  
 20 gctttccctt gcctgcctgt ctcttagtttc tctcacatcc cttttttttt tcctttctct  
 agccaccctg aagggtccct tcccaagccc tttagggacccg cagaggactt ggggaccaggc  
 aagcaacccc cagggcacga gaagagctct tgctgtctgc cctgcctcac cctgc(cc)ac  
 accaggccccg gtggcccccga gctgcataaa gtggaggcg aggaggaggc ggaggagggt  
 ggcaccatgg gcccgggcgg tgccctccat gcccggggga tgaagacact gctgcc atg  
 25 Met  
 1 347  
 gac agc ccg tgc cag ccg cag ccc cta agt cag gct ctc cct cag tta  
 Asp Ser Pro Cys Gln Pro Gln Pro Leu Ser Gln Ala Leu Pro Gln Leu  
 5 10 15  
 30 cca ggg tct tcg tca gag ccc ttg gag cct gag cct ggc cg<sup>g</sup> gcc agg  
 Pro Gly Ser Ser Ser Glu Pro Leu Glu Pro Glu Pro Gly Arg Ala Arg  
 20 25 30 395  
 atg gga gtg gag agt tac ctg ccc tgt ccc ctg ctc ccc tcc tac cac  
 Met Gly Val Glu Ser Tyr Leu Pro Cys Pro Leu Leu Pro Ser Tyr His  
 35 35 40 45 443  
 tgt cca gga gtg cct agt gag gcc tcg gca ggg agt ggg acc ccc aga  
 Cys Pro Gly Val Pro Ser Glu Ala Ser Ala Gly Ser Gly Thr Pro Arg  
 50 55 60 65 491  
 gcc aca gcc acc tct acc act gcc agc cct ctt cg<sup>g</sup> gac ggt ttt ggc  
 Ala Thr Ala Thr Ser Thr Ala Ser Pro Leu Arg Asp Gly Phe Gly  
 40 70 75 80 539  
 ggg cag gat ggt ggt gag ctg cg<sup>g</sup> ccg ctg cag agt gaa ggc gct gca  
 Gly Gln Asp Gly Gly Glu Leu Arg Pro Leu Gln Ser Glu Gly Ala Ala  
 85 90 95 587  
 45 gcg ctg gtc acc aag ggg tgc cag cga ttg gca gcc cag ggc gca cg<sup>g</sup>  
 Ala Leu Val Thr Lys Gly Cys Gln Arg Leu Ala Ala Gln Gly Ala Arg  
 100 105 110 635  
 cct gag gcc ccc aaa cg<sup>g</sup> aaa tgg gcc gag gat ggt ggg gat gcc cct  
 Pro Glu Ala Pro Lys Arg Lys Trp Ala Glu Asp Gly Gly Asp Ala Pro  
 115 120 125 683

	115	120	125	
	tca ccc agc aaa cgg ccc tgg gcc agg caa gag aac cag gag gca gag			731
	Ser Pro Ser Lys Arg Pro Trp Ala Arg Gln Glu Asn Gln Glu Ala Glu			
5	130 135 140 145			
	cgg gag ggt ggc atg agc tgc agc agt ggc agt ggt gag gcc			779
	Arg Glu Gly Gly Met Ser Cys Ser Ser Gly Ser Gly Glu Ala			
	150 155 160			
	agt gct ggg ctg atg gag gag gcg ctg ccc tct gcg ccc gag cgc ctg			827
	Ser Ala Gly Leu Met Glu Glu Ala Leu Pro Ser Ala Pro Glu Arg Leu			
10	165 170 175			
	gcc ctg gac tat atc gtg ccc tgc atg cgg tac tac ggc atc tgc gtc			875
	Ala Leu Asp Tyr Ile Val Pro Cys Met Arg Tyr Tyr Gly Ile Cys Val			
	180 185 190			
	aag gac agc ttc ctg ggg gca gca ctg ggc ggt cgc gtc ctg gcc gag			923
15	Lys Asp Ser Phe Leu Gly Ala Ala Leu Gly Gly Arg Val Leu Ala Glu			
	195 200 205			
	gtg gag gcc ctc aaa cgg ggt ggg cgc ctg cga gac ggg cag cta gtg			971
	Val Glu Ala Leu Lys Arg Gly Gly Arg Leu Arg Asp Gly Gln Leu Val			
	210 215 220 225			
20	agc cag agg gcg atc ccg ccg cgc agc atc cgt ggg gac cag att gcc			1019
	Ser Gln Arg Ala Ile Pro Pro Arg Ser Ile Arg Gly Asp Gln Ile Ala			
	230 235 240			
	tgg gtg gaa ggc cat gaa cca ggc tgt cga agc att ggt gcc ctc atg			1067
	Trp Val Glu Gly His Glu Pro Gly Cys Arg Ser Ile Gly Ala Leu Met			
25	245 250 255			
	gcc cat gtg gac gcc gtc atc cgc cac tgc gca ggg cgg ctg gcc agc			1115
	Ala His Val Asp Ala Val Ile Arg His Cys Ala Gly Arg Leu Gly Ser			
	260 265 270			
	tat gtc atc aac ggg cgc acc aag gcc atg gtg gcg tgt tac cca ggc			1163
30	Tyr Val Ile Asn Gly Arg Thr Lys Ala Met Val Ala Cys Tyr Pro Gly			
	275 280 285			
	aac ggg ctc ggg tac gta agg cac gtt gac aat ccc cac ggc gat ggg			1211
	Asn Gly Leu Gly Tyr Val Arg His Val Asp Asn Pro His Gly Asp Gly			
	290 295 300 305			
35	cgc tgc atc acc tgt atc tat tac ctg aat cag aac tgg gac gtt aag			1259
	Arg Cys Ile Thr Cys Ile Tyr Tyr Leu Asn Gln Asn Trp Asp Val Lys			
	310 315 320			
	gtg cat ggc ggc ctg ctg cag atc ttc cct gag ggc cgg ccc gtg gta			1307
	Val His Gly Leu Leu Gln Ile Phe Pro Glu Gly Arg Pro Val Val			
40	325 330 335			
	gcc aac atc gag cca ctc ttt gac cgg ttg ctc att ttc tgg tct gac			1355
	Ala Asn Ile Glu Pro Leu Phe Asp Arg Leu Leu Ile Phe Trp Ser Asp			
	340 345 350			
	cgg cgg aac ccc cac gag gtg aag cca gcc tat gcc acc agg tac gcc			1403
45	Arg Arg Asn Pro His Glu Val Lys Pro Ala Tyr Ala Thr Arg Tyr Ala			
	355 360 365			
	atc act gtc tgg tat ttt gat gcc aag gag cgg gca gca gcc aaa gac			1451
	Ile Thr Val Trp Tyr Phe Asp Ala Lys Glu Arg Ala Ala Lys Asp			
	370 375 380 385			

	aag tat cag cta gca tca gga cag aaa ggt gtc caa gta cct gta tca	1499
	Lys Tyr Gln Leu Ala Ser Gly Gln Lys Gly Val Gln Val Pro Val Ser	
	390 395 400	
5	cag ccg cct acg ccc acc tagtggccag tcccagagcc gcatggcaga	1547
	Gln Pro Pro Thr Pro Thr	
	405	
10	cagcttaaat gacttcagga gagccctggg cctgtgctgg ctgctccttc cctgccaccg ctgctgcttc tgactttgcc tctgtcctgc ctgggtgtgga gggctctgtc tggtgttag gaccaaggag gagaagagac ctttgcgtgcc ccatcatggg ggctgggggtt gtcacctgga	1607 1667 1727 1787 1847 1907 1967 2027 2087 2110
	cagggggcag cctgtggaggc caccgttacc aactgaagct gggggcctgg gtcctaccct gtctggcat gacccatta ggtatggaga gctggggagga ggcattgtca cttccccacca ggatgcagga cttggggttt aggtgagtca tggcctcttg ctggcaatgg ggtggggagga	
	gtaccccaa gtcctctcac tcctccagcc tggaatgtga agtactccc caacccctt ggccatggca ggcacccccc ggactgggct gccactgctt gggcagagta aaaggtgcca	
15	ggaggagcat gggtgtggaa gtccctgtcag ccaagaataaa aagtttacc tcagagctgc aaaaaaaaaaa aaaaaaaaaaaa aaa	
	<210> 2	
20	<211> 407	
	<212> PRT	
	<213> Homo sapiens	
	<400> 2	
25	Met Asp Ser Pro Cys Gln Pro Gln Pro Leu Ser Gln Ala Leu Pro Gln 1 5 10 15	
	Leu Pro Gly Ser Ser Ser Glu Pro Leu Glu Pro Glu Pro Gly Arg Ala 20 25 30	
	Arg Met Gly Val Glu Ser Tyr Leu Pro Cys Pro Leu Leu Pro Ser Tyr 35 40 45	
30	His Cys Pro Gly Val Pro Ser Glu Ala Ser Ala Gly Ser Gly Thr Pro 50 55 60	
	Arg Ala Thr Ala Thr Ser Thr Ala Ser Pro Leu Arg Asp Gly Phe 65 70 75 80	
	Gly Gly Gln Asp Gly Gly Glu Leu Arg Pro Leu Gln Ser Glu Gly Ala 85 90 95	
35	Ala Ala Leu Val Thr Lys Gly Cys Gln Arg Leu Ala Ala Gln Gly Ala 100 105 110	
	Arg Pro Glu Ala Pro Lys Arg Lys Trp Ala Glu Asp Gly Gly Asp Ala 115 120 125	
40	Pro Ser Pro Ser Lys Arg Pro Trp Ala Arg Gln Glu Asn Gln Glu Ala 130 135 140	
	Glu Arg Glu Gly Gly Met Ser Cys Ser Cys Ser Ser Gly Ser Gly Glu 145 150 155 160	
	Ala Ser Ala Gly Leu Met Glu Glu Ala Leu Pro Ser Ala Pro Glu Arg 165 170 175	
45	Leu Ala Leu Asp Tyr Ile Val Pro Cys Met Arg Tyr Tyr Gly Ile Cys 180 185 190	
	Val Lys Asp Ser Phe Leu Gly Ala Ala Leu Gly Gly Arg Val Leu Ala 195 200 205	

	Glu Val Glu Ala Leu Lys Arg Gly Gly Arg Leu Arg Asp Gly Gln Leu	
	210 215 220	
	Val Ser Gln Arg Ala Ile Pro Pro Arg Ser Ile Arg Gly Asp Gln Ile	
	225 230 235 240	
5	Ala Trp Val Glu Gly His Glu Pro Gly Cys Arg Ser Ile Gly Ala Leu	
	245 250 255	
	Met Ala His Val Asp Ala Val Ile Arg His Cys Ala Gly Arg Leu Gly	
	260 265 270	
10	Ser Tyr Val Ile Asn Gly Arg Thr Lys Ala Met Val Ala Cys Tyr Pro	
	275 280 285	
	Gly Asn Gly Leu Gly Tyr Val Arg His Val Asp Asn Pro His Gly Asp	
	290 295 300	
	Gly Arg Cys Ile Thr Cys Ile Tyr Tyr Leu Asn Gln Asn Trp Asp Val	
	305 310 315 320	
15	Lys Val His Gly Gly Leu Leu Gln Ile Phe Pro Glu Gly Arg Pro Val	
	325 330 335	
	Val Ala Asn Ile Glu Pro Leu Phe Asp Arg Leu Leu Ile Phe Trp Ser	
	340 345 350	
20	Asp Arg Arg Asn Pro His Glu Val Lys Pro Ala Tyr Ala Thr Arg Tyr	
	355 360 365	
	Ala Ile Thr Val Trp Tyr Phe Asp Ala Lys Glu Arg Ala Ala Ala Lys	
	370 375 380	
	Asp Lys Tyr Gln Leu Ala Ser Gly Gln Lys Gly Val Gln Val Pro Val	
	385 390 395 400	
25	Ser Gln Pro Pro Thr Pro Thr	
	405	
	<210> 3	
	<211> 5163	
30	<212> DNA	
	<213> Homo sapiens	
	<220>	
35	<221> CDS	
	<222> (3157)..(4434)	
	<223>	
	<400> 3	
40	ttagggcag aaaaacattt gtaataatta atggctttga gagacacaag gctttgtttg ccccagagta tttagtaacc cacctagtgc tcctaattcat acaatattaa ggattgggag ggacattcat tgcctcaactc tctatttgtt tcaccttctg taaaatttgtt agaataatag tacccacttc atagcattgt atgatgattaa aattggtaa tatttttaaa atgcttagaa cacagattgg gcacataaca gcaaggcacca catgtgttta taagataaat tcctttgtgt	60 120 180 240 300
45	tgccttcgt taaagttaa ataagtaaat aaataaataa atacttgcat gacattttga agtctctcta taacatctga gtaagtggcg gctgcgacaa tgctactgga gttccagaat cgtgttggtg acaagattgt tcaccagcat atggtgtggtaaaaactcac taatttgaa ttagttcaga ttattaagcc tgaataggtaaaaatccctga aatcaaggat ctttggaaact atttqaaatc agtattttat atttcctgt tgtattcatt aaagtgttgc aagtgttca	360 420 480 540 600



	Leu Cys Gly Lys Met Glu Asn Leu Leu Arg Cys Ser Arg Cys Arg Ser				
	25	30	35		
	tcc ttc tac tgc tgc aag gag cac cag cgt cag gac tgg aag aag cac			3318	
	Ser Phe Tyr Cys Cys Lys Glu His Gln Arg Gln Asp Trp Lys Lys His				
5	40	45	50		
	aag ctc gtg tgc cag ggc agc gag ggc gcc ctc ggc cac gga gtg ggc			3366	
	Lys Leu Val Cys Gln Gly Ser Glu Gly Ala Leu Gly His Gly Val Gly				
	55	60	65	70	
10	cca cac cag cat tcc ggc ccc gcg ccg gct gca gtg ccg ccg ccc			3414	
	Pro His Gln His Ser Gly Pro Ala Pro Pro Ala Ala Val Pro Pro Pro				
	75	80	85		
	agg gcc ggg gcc cggttccggccagg aag gca gcg gcg cgg gac aac			3462	
	Arg Ala Gly Ala Arg Glu Pro Arg Lys Ala Ala Arg Arg Asp Asn				
	90	95	100		
15	gcc tcc ggg gac gcg gcc aag gga aaa gta aag gcc aag ccc ccg gcc			3510	
	Ala Ser Gly Asp Ala Ala Lys Gly Lys Val Lys Ala Lys Pro Pro Ala				
	105	110	115		
	gac cca gcg gcg gcc gcg tcg ccg tgt cgt gcg gcc ggc ggc cag			3558	
	Asp Pro Ala Ala Ala Ser Pro Cys Arg Ala Ala Gly Gly Gln				
20	120	125	130		
	ggc tcg gcg gtg gct gcc gaa gcc gag ccc ggc aag gag gag ccg ccg			3606	
	Gly Ser Ala Val Ala Ala Glu Ala Glu Pro Gly Lys Glu Glu Pro Pro				
	135	140	145	150	
25	gcc cgc tca tcg ctg ttc cag gag aag gcg aac ctg tac ccc cca agc			3654	
	Ala Arg Ser Ser Leu Phe Gln Glu Lys Ala Asn Leu Tyr Pro Pro Ser				
	155	160	165		
	aac acg ccc ggg gat gcg ctg agc ccc ggc ggc ctg cgg ccc aac			3702	
	Asn Thr Pro Gly Asp Ala Leu Ser Pro Gly Gly Leu Arg Pro Asn				
	170	175	180		
30	ggg cag acg aag ccc ctg ccg gcg ctg aag ctg gcg ctc gag tac atc			3750	
	Gly Gln Thr Lys Pro Leu Pro Ala Leu Lys Leu Ala Leu Glu Tyr Ile				
	185	190	195		
	gtg ccg tgc atg aac aag cac ggc atc tgt gtg gac gac ttc ctc			3798	
	Val Pro Cys Met Asn Lys His Gly Ile Cys Val Val Asp Asp Phe Leu				
35	200	205	210		
	ggc aag gag acc gga cag cag atc ggc gac gag gtg cgc gcc ctg cac			3846	
	Gly Lys Glu Thr Gly Gln Gln Ile Gly Asp Glu Val Arg Ala Leu His				
	215	220	225	230	
40	gac acc ggg aag ttc acg gac ggg cag ctg gtc agc cag aag agt gac			3894	
	Asp Thr Gly Lys Phe Thr Asp Gly Gln Leu Val Ser Gln Lys Ser Asp				
	235	240	245		
	tcg tcc aag gac atc cga ggc gat aag atc acc tgg atc gag ggc aag			3942	
	Ser Ser Lys Asp Ile Arg Gly Asp Lys Ile Thr Trp Ile Glu Gly Lys				
	250	255	260		
45	gag ccc ggc tgc gaa acc att ggg ctg ctc atg agc agc atg gac gac			3990	
	Glu Pro Gly Cys Glu Thr Ile Gly Leu Leu Met Ser Ser Met Asp Asp				
	265	270	275		
	ctg ata cgc cac tgt aac ggg aag ctg ggc agc tac aaa atc aat ggc			4038	
	Leu Ile Arg His Cys Asn Gly Lys Leu Gly Ser Tyr Lys Ile Asn Gly				

	280	285	290	
	cgg acg aaa gcc atg gtt gct tgt tat ccg ggc aat gga acg ggt tat			
	Arg Thr Lys Ala Met Val Ala Cys Tyr Pro Gly Asn Gly Thr Gly Tyr			
5	295	300	305	310
	gta cgt cat gtt gat aat cca aat gga gat gga aga tgt gtg aca tgt			
	Val Arg His Val Asp Asn Pro Asn Gly Asp Gly Arg Cys Val Thr Cys			
	315	320	325	
	ata tat tat ctt aat aaa gac tgg gat gcc aag gta agt gga ggt ata			4182
	Ile Tyr Tyr Leu Asn Lys Asp Trp Asp Ala Lys Val Ser Gly Gly Ile			
10	330	335	340	
	ctt cga att ttt cca gaa ggc aaa gcc cag ttt gct gac att gaa ccc			4230
	Leu Arg Ile Phe Pro Glu Gly Lys Ala Gln Phe Ala Asp Ile Glu Pro			
	345	350	355	
15	aaa ttt gat aga ctg ctg ttt ttc tgg tct gac cgt cgc aac cct cat			4278
	Lys Phe Asp Arg Leu Leu Phe Phe Trp Ser Asp Arg Arg Asn Pro His			
	360	365	370	
	gaa gta caa cca gca tat gct aca agg tac gca ata act gtt tgg tat			
	Glu Val Gln Pro Ala Tyr Ala Thr Arg Tyr Ala Ile Thr Val Trp Tyr			
	375	380	385	390
20	ttt gat gca gat gag aga gca cga gct aaa gta aaa tat cta aca ggt			4326
	Phe Asp Ala Asp Glu Arg Ala Arg Ala Lys Val Lys Tyr Leu Thr Gly			
	395	400	405	
	gaa aaa ggt gtg agg gtt gaa ctc aat aaa cct tca gat tcg gtc ggt			4422
	Glu Lys Gly Val Arg Val Glu Leu Asn Lys Pro Ser Asp Ser Val Gly			
25	410	415	420	
	aaa gac gtc ttc tagaggccttt gatccagcaa taccggactt cacctacaat			4474
	Lys Asp Val Phe			
	425			
	attgttaact atttgttaac ttgtgaatac gaataaatgg gataaaagaaa aatagacaac			4534
30	cagttcgcatttataataagg aaacagaaaac aactttttgt ttgcatacaa acagaagatt			4594
	ttgactgttgactttgtt ctgcataatgc aacttcaaattt ctgtgattgc ttacaggagg			4654
	aagataagct actaatttggaa aatgggtttt acatctggat atggaaataag tgccctgtgt			4714
	agaattttttt tcatttttat attttgcag atctgttatac tagttagttt cattttcatct			4774
	ctccctttttt tataatcaagt ttgaattttgg gataattttt ctatattagg tacaattttat			4834
35	ctaaactgaa ttgagaaaaa attacagtat tatttcctcaa aataacatca atctattttt			4894
	gttaaaacctgt tcataactatt aaattttgcc cttaaaagacc tcttaataat gattgttgc			4954
	agtgactgtat gattaattttt attttactta aaataagaaaa aggaggactt taattacaac			5014
	tgaaaaatca gatttttttgc agtccttcc ttacactaat ttgaacttctt aaagattgt			5074
	gctttttttt tgacattgtc aataacgaaa cctaattgtaa acacagtcac cattttactac			5134
40	caataactttt tagttaatgt tttacaagg			5163
	<210> 4			
	<211> 426			
	<212> PRT			
45	<213> Homo sapiens			
	<400> 4			

Met Ala Asn Asp Ser Gly Gly Pro Gly Gly Pro Ser Pro Ser Glu Arg  
 1                   5                   10                   15

Asp Arg Gln Tyr Cys Glu Leu Cys Gly Lys Met Glu Asn Leu Leu Arg  
                  20                 25                 30  
 Cys Ser Arg Cys Arg Ser Ser Phe Tyr Cys Cys Lys Glu His Gln Arg  
                  35                 40                 45  
 5   Gln Asp Trp Lys His Lys Leu Val Cys Gln Gly Ser Glu Gly Ala  
       50                 55                 60  
 Leu Gly His Gly Val Gly Pro His Gln His Ser Gly Pro Ala Pro Pro  
       65                 70                 75                 80  
 Ala Ala Val Pro Pro Pro Arg Ala Gly Ala Arg Glu Pro Arg Lys Ala  
       85                 90                 95  
 10   Ala Ala Arg Arg Asp Asn Ala Ser Gly Asp Ala Ala Lys Gly Lys Val  
       100                105                110  
 Lys Ala Lys Pro Pro Ala Asp Pro Ala Ala Ala Ser Pro Cys Arg  
       115                120                125  
 15   Ala Ala Ala Gly Gly Gln Gly Ser Ala Val Ala Ala Glu Ala Glu Pro  
       130                135                140  
 Gly Lys Glu Glu Pro Pro Ala Arg Ser Ser Leu Phe Gln Glu Lys Ala  
       145                150                155                160  
 Asn Leu Tyr Pro Pro Ser Asn Thr Pro Gly Asp Ala Leu Ser Pro Gly  
       165                170                175  
 20   Gly Gly Leu Arg Pro Asn Gly Gln Thr Lys Pro Leu Pro Ala Leu Lys  
       180                185                190  
 Leu Ala Leu Glu Tyr Ile Val Pro Cys Met Asn Lys His Gly Ile Cys  
       195                200                205  
 25   Val Val Asp Asp Phe Leu Gly Lys Glu Thr Gly Gln Gln Ile Gly Asp  
       210                215                220  
 Glu Val Arg Ala Leu His Asp Thr Gly Lys Phe Thr Asp Gly Gln Leu  
       225                230                235                240  
 Val Ser Gln Lys Ser Asp Ser Ser Lys Asp Ile Arg Gly Asp Lys Ile  
       245                250                255  
 30   Thr Trp Ile Glu Gly Lys Glu Pro Gly Cys Glu Thr Ile Gly Leu Leu  
       260                265                270  
 Met Ser Ser Met Asp Asp Leu Ile Arg His Cys Asn Gly Lys Leu Gly  
       275                280                285  
 35   Ser Tyr Lys Ile Asn Gly Arg Thr Lys Ala Met Val Ala Cys Tyr Pro  
       290                295                300  
 Gly Asn Gly Thr Gly Tyr Val Arg His Val Asp Asn Pro Asn Gly Asp  
       305                310                315                320  
 Gly Arg Cys Val Thr Cys Ile Tyr Tyr Leu Asn Lys Asp Trp Asp Ala  
       325                330                335  
 40   Lys Val Ser Gly Gly Ile Leu Arg Ile Phe Pro Glu Gly Lys Ala Gln  
       340                345                350  
 Phe Ala Asp Ile Glu Pro Lys Phe Asp Arg Leu Leu Phe Phe Trp Ser  
       355                360                365  
 45   Asp Arg Arg Asn Pro His Glu Val Gln Pro Ala Tyr Ala Thr Arg Tyr  
       370                375                380  
 Ala Ile Thr Val Trp Tyr Phe Asp Ala Asp Glu Arg Ala Arg Ala Lys  
       385                390                395                400  
 Val Lys Tyr Leu Thr Gly Glu Lys Gly Val Arg Val Glu Leu Asn Lys

	405	410	415	
	Pro Ser Asp Ser Val Gly Lys Asp Val Phe			
	420	425		
5	<210> 5			
	<211> 2770			
	<212> DNA			
	<213> Homo sapiens			
	<220>			
10	<221> CDS			
	<222> (327)..(1043)			
	<223>			
	 5			
15	gagtctggcc gcagtcgcgg cagtggtggc ttccccatccc caaaaaggcgcc cctccgactc cttgcgccgc actgctcgcc gggccagtcgg ggaaacgggt cgtggagctc cgaccactc ccgctggttc ccgaaggcag atcccttetc ccgagagttt cgagaaaactt tcccttgcc ccgacgctgc agcggctcgg gtaccgtggc agccgcagggt ttctgaaccc cgggcccacgc tccccgcgcc tcggcttcgc gctcgtgttag atcggtccct ctctggttgc acgctgggga	60 120 180 240 300 353		
20	tcccgaccc cgattctgcg ggcgag atg ccc ctg gga cac atc atg agg ctg Met Pro Leu Gly His Ile Met Arg Leu 1 5			
	gac ctg gag aaa att gcc ctg gag tac atc gtg ccc tgt ctg cac gag Asp Leu Glu Lys Ile Ala Leu Glu Tyr Ile Val Pro Cys Leu His Glu	401		
25	10 15 20 25			
	gtg ggc ttc tgc tac ctg gac aac ttc ctg ggc gag gtg gtg ggc gac Val Gly Phe Cys Tyr Leu Asp Asn Phe Leu Gly Glu Val Val Gly Asp 30 35 40	449		
30	tgc gtc ctg gag cgc gtc aag cag ctg cac tgc acc ggg gcc ctg cgg Cys Val Leu Glu Arg Val Lys Gln Leu His Cys Thr Gly Ala Leu Arg 45 50 55	497		
	gac ggc cag ctg gcg ggg ccg cgc gcc ggc gtc tcc aag cga cac ctg Asp Gly Gln Leu Ala Gly Pro Arg Ala Gly Val Ser Lys Arg His Leu 60 65 70	545		
35	cgg ggc gac cag atc acg tgg atc ggg ggc aac gag gag ggc tgc gag Arg Gly Asp Gln Ile Thr Trp Ile Gly Gly Asn Glu Glu Gly Cys Glu 75 80 85	593		
	gcc atc agc ttc ctc ctg tcc ctc atc gac agg ctg gtc ctc tac tgc Ala Ile Ser Phe Leu Leu Ser Leu Ile Asp Arg Leu Val Leu Tyr Cys	641		
40	90 95 100 105			
	ggg agc cgg ctg ggc aaa tac tac gtc aag gag agg tct aag gca atg Gly Ser Arg Leu Gly Lys Tyr Tyr Val Lys Glu Arg Ser Lys Ala Met 110 115 120	689		
45	gtg gct tgc tat ccg gga aat gga aca ggt tat gtt cgc cac gtg gac Val Ala Cys Tyr Pro Gly Asn Gly Thr Gly Tyr Val Arg His Val Asp 125 130 135	737		
	aac ccc aac ggt gat ggt cgc tgc atc acc tgc atc tac tat ctg aac Asn Pro Asn Gly Asp Gly Arg Cys Ile Thr Cys Ile Tyr Tyr Leu Asn 140 145 150	785		

	aag aat tgg gat gcc aag cta cat ggt ggg atc ctg cgg ata ttt cca	833
	Lys Asn Trp Asp Ala Lys Leu His Gly Ile Leu Arg Ile Phe Pro	
	155 160 165	
5	gag ggg aaa tca ttc ata gca gat gtg gag ccc att ttt gac aga ctc	881
	Glu Gly Lys Ser Phe Ile Ala Asp Val Glu Pro Ile Phe Asp Arg Leu	
	170 175 180 185	
	ctg ttc ttc tgg tca gat cgt agg aac cca cac gaa gtg cag ccc tct	929
	Leu Phe Phe Trp Ser Asp Arg Arg Asn Pro His Glu Val Gln Pro Ser	
	190 195 200	
10	tac gca acc aga tat gct atg act gtc tgg tac ttt gat gct gaa gaa	977
	Tyr Ala Thr Arg Tyr Ala Met Thr Val Trp Tyr Phe Asp Ala Glu Glu	
	205 210 215	
	agg gca gaa gcc aaa aag aaa ttc agg aat tta act agg aaa act gaa	1025
	Arg Ala Glu Ala Lys Lys Phe Arg Asn Leu Thr Arg Lys Thr Glu	
15	220 225 230	
	tct gcc ctc act gaa gac tgaccgtgct ctgaaatctg ctggccttgc	1073
	Ser Ala Leu Thr Glu Asp	
	235	
20	tcatTTtagt aacggttccct gaattctctt aaattctttg agatccaaag atggccttctt	1133
	cagtgacaac aatcicccctg ctacttcttg catccttcac atcccctgtct tgtgtgtgg	1193
	acttcatgtt ttcttgccaa gactgtgttg atcttcagat actctctttg ccagatgaag	1253
	ttatTTgcta actccagaaa ttccctgcaga catcctactc ggccagcggt ttacctgata	1313
	gattcggtaa tactatcaag agaagagcct aggagcacag cgagggaatg aaccttactt	1373
25	gcactttatg tatacttcct gatTTgaaag gaggaggTTT gaaaaagaaaa aatgggagg	1433
	ggttagatgcc acagagaggc atcacggaaag ccttaacagc aggaaacaga gaaatttgtg	1493
	tcatctgaac aatTTccaga ttttcttaat ccagggtgt tggtgtttct ggagaattat	1553
	cacaacctaa tgacattaat acctctagaa agggctgtcg tcatagtgaa caatTTtataa	1613
	gtgtcccatg gggcagacac tccttttttc ccagtccgtc aacctggatt ttctgcctca	1673
	gctccatttt gctgaaaata atgactttct gaataaaagat ggcaacacaa ttttttctcc	1733
30	atTTtcagtt cttacctggg aacctaattc cccagaagct aaaaaactag acattagtt	1793
	ttttggTTgc ttTgTTggaa tggaaTTaa atTTaaatga aaggaaaaat atatccctgg	1853
	tagTTTGTG ttaaccactg ataactgtgg aaagagctag gtctactgtatacaataaaa	1913
	catgtgtca tcttgaacaa ttTgagaggg gaggtggagt tggaaatgtg ggtgttcctg	1973
	tttttttttt tttttttttt ttttttttagt tttcctttt aatgagctca ccTTtaaca	2033
35	aaaaaaaaagc agggTgatgt atTTaaaaaa aggaagtggaa aaaaaaaaaa tctcaaagct	2093
	atTTgagttc tcgtctgtcc ctagcagtct ttcttcagct cactggctc tctagatcca	2153
	ctgtggTTgg cagtagtgc accatcatgg aacttgcgt aactgtggaa gtttctactc	2213
	ctgcagtagc cacagatcgc actgcctcaa taactggta ttgagcacgt atTTTgcAAA	2273
	agctactttt cctagTTTC agtattactt tcataTTTTT aaaaatccTT taatTTcttg	2333
40	ctTgaaaatc ccatgaacat taaagagcCA gaaatTTTTT cctttgttat gtacggat	2393
	atatatatat atagTcttcc aagatagaag ttTactTTT cctttctgg tttTggaaaa	2453
	tttccagata agacatgtca ccattaattc tcaacgactg ctctatTTt tgTacggta	2513
	atagTTatca ccttctaaat tactatgtaa ttTactact tattatgttt attgtcttg	2573
	atccTTtctc tggagtgtaa gcacaatgaa gacaggaatt ttgtatTTT ttaaccaatg	2633
45	caacatactc tcagcaccta aaatagtGCC gggAACATAG taagggctca gtaaaatactt	2693
	gttgaataaa ctcagtctcc tacattagca ttctaaaaaa aaaaaaaaaa aaaaaaaaaa	2753
	aaaaaaaaaaaa aaaaaaag	2770

<211> 239  
<212> PRT  
<213> Homo sapiens  
<400> 6

5 Met Pro Leu Gly His Ile Met Arg Leu Asp Leu Glu Lys Ile Ala Leu  
1 5 10 15  
Glu Tyr Ile Val Pro Cys Leu His Glu Val Gly Phe Cys Tyr Leu Asp  
20 25 30  
10 Asn Phe Leu Gly Glu Val Val Gly Asp Cys Val Leu Glu Arg Val Lys  
35 40 45  
Gln Leu His Cys Thr Gly Ala Leu Arg Asp Gly Gln Leu Ala Gly Pro  
50 55 60  
Arg Ala Gly Val Ser Lys Arg His Leu Arg Gly Asp Gln Ile Thr Trp  
65 70 75 80  
15 Ile Gly Gly Asn Glu Glu Gly Cys Glu Ala Ile Ser Phe Leu Leu Ser  
85 90 95  
Leu Ile Asp Arg Leu Val Leu Tyr Cys Gly Ser Arg Leu Gly Lys Tyr  
100 105 110  
20 Tyr Val Lys Glu Arg Ser Lys Ala Met Val Ala Cys Tyr Pro Gly Asn  
115 120 125  
Gly Thr Gly Tyr Val Arg His Val Asp Asn Pro Asn Gly Asp Gly Arg  
130 135 140  
25 Cys Ile Thr Cys Ile Tyr Tyr Leu Asn Lys Asn Trp Asp Ala Lys Leu  
145 150 155 160  
His Gly Gly Ile Leu Arg Ile Phe Pro Glu Gly Lys Ser Phe Ile Ala  
165 170 175  
Asp Val Glu Pro Ile Phe Asp Arg Leu Leu Phe Phe Trp Ser Asp Arg  
180 185 190  
30 Arg Asn Pro His Glu Val Gln Pro Ser Tyr Ala Thr Arg Tyr Ala Met  
195 200 205  
Thr Val Trp Tyr Phe Asp Ala Glu Glu Arg Ala Glu Ala Lys Lys Lys  
210 215 220  
35 Phe Arg Asn Leu Thr Arg Lys Thr Glu Ser Ala Leu Thr Glu Asp  
225 230 235